#### SEQUENCE LISTING

- (1) GENERAL INFORMATION:
- (i) APPLICANT: FIDDES, J.C. ABRAHAM, J.D.
- (ii) TITLE OF INVENTION: HUMAN BASIC FIBROBLAST GROWTH FACTOR ANALOG
- (iii) NUMBER OF SEQUENCES: 69
- (iv) CORRESPONDENCE ADDRESS:
  - (A) ADDRESSEE: MORRISON & FOERSTER
  - (B) STREET: 755 PAGE MILL ROAD
  - (C) CITY: Palo Alto
  - (D) STATE: CA
  - (E) COUNTRY: USA
  - (F) ZIP: 94304-1018
- (v) COMPUTER READABLE FORM:
  - (A) MEDIUM TYPE: Diskette
  - (B) COMPUTER: IBM Compatible
  - (C) OPERATING SYSTEM: Windows
  - (D) SOFTWARE: FastSEQ for Windows Version 2.0b
- (vi) CURRENT APPLICATION DATA:
  - (A) APPLICATION NUMBER:
  - (B) FILING DATE:
- (C) CLASSIFICATION:
- (vii) PRIOR APPLICATION DATA:
  - (A) APPLICATION NUMBER:
  - (B) FILING DATE:
- (viii) ATTORNEY/AGENT INFORMATION:
  - (A) NAME: Lehnhardt, Susan K
  - (B) REGISTRATION NUMBER: 33,943
  - (C) REFERENCE/DOCKET NUMBER: 2190 \(\daggerq -20089.10\)
- (ix) TELECOMMUNICATION INFORMATION:
  - (A) TELEPHONE: 650-813-5600
  - (B) TELEFAX: 650-494-0792
  - (C) TELEX: 706141
  - (2) INFORMATION FOR SEQ ID NO:1:
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 1969 base pairs
  - (B) TYPE: nucleic acid
  - (C) STRANDEDNESS: double
  - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: cDNA

# (ix) FEATURE:

- (A) NAME/KEY: Coding Sequence (B) LOCATION: 970...1434
- (D) OTHER INFORMATION:

# (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

		( <b>X</b>	1) S	FOOF	NCE	DESC	RIPT	TON:	SEQ	עד	NO: I	•					
	TAAA GCTC TCCG AGTT CAGG CCCG GCAG AAGA CGGG CCCA AGAA CGGG GGCC	GTTT TGAC TTTTT GAGT GAGT GAAT GAGGG AAGC ATCTG AGAGC CGGGG	AT GA A CA	CCCC GACT ACGC GGGG AAAG CCGAA GCGG GCGC ACC ACCC ACCC ACCC ACCC ACCC ACCC ACCC ACCC ACCC ACCC ACCC ACCC ACC ACCC ACCC ACCC ACCC ACCC ACCC ACCC ACCC ACCC ACCC ACCC ACCC ACC ACCC ACCC ACCC ACCC ACCC ACCC ACCC ACCC ACCC ACCC ACCC ACCC ACC ACCC ACCC ACCC ACCC ACCC ACCC ACCC ACCC ACCC ACCC ACCC ACCC ACC	ACTT CFTC CTAGC CGGT ACTG ACTG ACTG CCAGG	G TA C TT C GG G GG G GG G GG G GG G GG G GG	GCCC GGAT GAAA GCACG GCGG GCGCG GCGGG CCGGG CCGGG	TCGT TGCA AAAT AAAA GGGT CCGC GCCC GGGA TGCC GCCGC TGGG GCCG GGCCG GGCCG GGCCG	CTT ACT GGG GCC GGC CGG CGG CGG GGC GGC GGC G	TTGG GGAG CCGC TTTTT CGCG AAAA TTGGT CTCCG CCCGG	TGA TAC AAA AGT GGG GCGC GCGG GCGG GCCC GCGG	TTTA TTTG GTTG GGAT CCCC TCGC GAGC TGTG ACGG ACG	GAGA GGGTT AGAA AAGG CGCCC GGGGG CGGCG CCCAG GCCGA CCCAG CCCAG CCCAG CCCAG CCCAG CCCAG CCCAG CCCAG CCCAG	TT T GG A AG C GG C CC T TT G GG C GG C GG C GG C G	TCAA AACG ACTT CTAA CGGTG CGCC CGCGC CGCGC CGCGC CGCGC CCCGC CCCGCC CCCGCC CCCGCC CCCGCC CCCGCCC CCCGCCC CCCGCCCCCC	AATTT AGCCT GCTTC TTAAA ACGTG GAGCC CTCTCC GTTGTG GGGGCC ATGTAG AGGCAA AGGCAC GCGGAC GCCGGG GGCCCC CC GAG TO Glu	60 120 180 240 300 360 420 480 540 600 660 720 780 840 900 960 1011
	GAT Asp 15	GGC Gly	GGC Gly	AGC Ser	GGC Gly	GCC Ala 20	TTC Phe	CCG Pro	CCC Pro	GGC Gly	CAC His 25	TTC Phe	AAG Lys	GAC Asp	CCC Pro	AAG Lys 30	1059
	CGG Arg	CTG Leu	TAC Tyr	TGC Cys	AAA Lys 35	AAC Asn	GGG Gly	ĠGC G <b>l</b> y	TTC Phe	TTC Phe 40	CTG Leu	CGC Arg	ATC Ile	CAC His	CCC Pro 45	GAC Asp	1107
)	GGC Gly	CGA Arg	GTT Val	GAC Asp 50	GGG Gly	GTC Val	CGG Arg	GAG\ Glu	AAG Lys \$5	AGC Ser	GAC Asp	CCT Pro	CAC His	ATC Ile 60	AAG Lys	CTA Leu	1155
	CAA Gln	CTT Leu	CAA Gln 65	GCA Ala	GAA Glu	GAG Glu	AGA Arg	GGA Gly 70	G <b>T</b> T Val	GTG Val	TCT Ser	ATC Ile	AAA Lys 75	GGA Gly	GTG Val	TGT Cys	1203
	GCT Ala	AAC Asn 80	CGT Arg	TAC Tyr	CTG Leu	GCT Ala	ATG Met 85	AAG Lys	GAA Glu	GAT Asp	GGA Gly	AGA Arg 90	TTA Leu	CTG Leu	GCT Ala	TCT Ser	1251
	AAA Lys 95	TGT Cys	GTT Val	ACG Thr	GAT Asp	GAG Glu 100	TG <b>T</b> Cys	TTC Phe	TTT Phe	TTT Phe	GAA Glu 105	CGA Arg	TTG Leu	GAA Glu	TCT Ser	AAT Asn 110	1299
	AAC Asn	TAC Tyr	AAT Asn	ACT Thr	TAC Tyr	CGG Arg	TCA Ser	AGG Arg	AAA Lys	TAC Tyr 120	ACC Thr	AGT Ser	TGG Trp	TAT Tyr	GTG Val 125	GCA Ala	1347

			ACT That 130													1395
			ATA Ile										TGA'	TTTT	AAT GG	1446
CCAC	CATC	ΓAA '	TCTC	ATTT	CA C	ATGA	AAGA!	A GA	AGTA'	TATT	TTA	GAAA'	rrr ·	GTTA	ATGAGA	1506
															TTATCO	
															TGTATA	
TTC	rccci	TTT '	TATA	гтфси	AT C	TGCT	GTTA(	CC	AGTG	AAGC	TTA	CCTA	GAG	CAAT	GATCTT	1686
TTT	CACG	CAT '	TTGC	rtta:	TT C	GAAA	AGAG	CT'	TTTA	TAA	GTG	CATG	TTT .	AGAA	AACAAA	. 1746

GTAAAAGAAA ATAAATGTGT ATAGCTCAGT TTGGATAATT GGTCAAACAA TTTTTTATCC 1566
AGTAGTAAAA TATGTAACCA TGCCCAGTAA AGAAAAATAA CAAAAGTTGT AAAATGTATA 1626
TTCTCCCTTT TATATTCAT CTGCTGTTAC CCAGTGAAGC TTACCTAGAG CAATGATCTT 1686
TTTCACGCAT TTGCTTTATT CGAAAAGAGG CTTTTAAAAT GTGCATGTTT AGAAAACAAA 1746
AATGTCCACT ATTTCTTATG TCATTCGTTA GTCTACATGT TTCTAAACAT ATAAATGTGA 1806
ATTTAATCAA TTCCTTTCAT AGTTTTATAA TTCTCTGGCA GTTCCTTATG ATAGAGTTTA 1926
TAAAACAGTC CTGTGTAAAC TGCTGGAAGT TCTTCCGGAA TTC 1969

# (2) INFORMATION FOR SEQ ID NO:2:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 155 amino acids
  - (B) TYPE: amind acid
  - (C) STRANDEDNESS: single
  - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE:\protein
- (v) FRAGMENT TYPE: internal

## (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

Met Ala Ala Gly Ser Ile Thr Thr Leu Pro Ala Leu Pro Glu Asp Gly 10 Gly Ser Gly Ala Phe Pro Pro Gly His Phe Lys Asp Pro Lys Arg Leu 25 Tyr Cys Lys Asn Gly Gly Phe Phe Leu Arg Ile His Pro Asp Gly Arg Val Asp Gly Val Arg Glu Lys Sen Asp Pro His Ile Lys Leu Gln Leu Gln Ala Glu Glu Arg Gly Val Val\Ser Ile Lys Gly Val Cys Ala Asn 70 Arg Tyr Leu Ala Met Lys Glu Asp Gly Arg Leu Leu Ala Ser Lys Cys 90 85 Val Thr Asp Glu Cys Phe Phe Phe Glu Arg Leu Glu Ser Asn Asn Tyr 1**(**5 100 Asn Thr Tyr Arg Ser Arg Lys Tyr Thr Ser Trp Tyr Val Ala Leu Lys 120 Arg Thr Gly Gln Tyr Lys Leu Gly Set Lys Thr Gly Pro Gly Gln Lys 135 Ala Ile Leu Phe Leu Pro Met Ser Ala Lys Ser 150 145

### (2) INFORMATION FOR SEQ ID NO:3:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 638 base pairs
  - (B) TYPE: nucleic acid

- (C) STRANDEDNESS: double (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: cDNA
- (ix) FEATURE:
  - (A) NAME/KEY: Coding Sequence
  - (B) LOCATION: 91...555
  - (D) OTHER INFORMATION:
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

		( X	(1) S	EQUE	NCE	DESC	RIPT	TON:	SEC	י דט	NO:3						
	TGCA GAAA	ATTTT AGCGC	GT G	GCCTT CAAGC	TGCT CAGCA	'G GA LG CT	AGAA 'GCTG	CCGA AGCC	ra :	G GC t Al	T GA	AA GO	G G	AA AT .u Il	C AC	GTCTT C ACC ir Thr	60 114
	TTC Phe	ACA Thr 10	GCC Ala	CTG Leu	ACC	GAG Glu	AAG Lys 15	TTT Phe	AAT Asn	CTG Leu	CCT Pro	CCA Pro 20	GGG Gly	AAT Asn	TAC Tyr	AAG Lys	162
	AAG Lys 25	CCC Pro	AAA Lys	CTC Leu	CTC Leu	TAC Tyr 30	TGT Cys	AGC Ser	AAC Asn	GGG Gly	GGC Gly 35	CAC His	TTC Phe	CTG Leu	AGG Arg	ATC Ile 40	210
	CTT Leu	CCG Pro	GAT Asp	GGC Gly	ACA Thr 45	GTG\ Val	GAT Asp	GGG Gly	ACA Thr	AGG Arg 50	GAC Asp	AGG Arg	AGC Ser	GAC Asp	CAG Gln 55	CAC His	258
	ATT Ile	CAG Gln	CTG Leu	CAG Gln 60	CTC Leu	AGT Ser	GCG Ala	GAA Glu	AGC Ser 65	GTG Val	GGG Gly	GAG Glu	GTG Val	TAT Tyr 70	ATA Ile	AAG Lys	306
	AGT Ser	ACC Thr	GAG Glu 75	ACT Thr	GGC Gly	CAG Gln	TAC	TTG Leu 80	GCC Ala	ATG Met	GAC Asp	ACC Thr	GAC Asp 85	GGG Gly	CTT Leu	TTA Leu	354
1	TAC Tyr	GGC Gly 90	TCA Ser	CAG Gln	ACA Thr	CCA Pro	AAT Asn 95	GAG Glu	GAA Glu	TGT Cys	TTG Leu	TTC Phe 100	CTG Leu	GAA Glu	AGG Arg	CTG Leu	402
,	GAG Glu 105	GAG Glu	AAC Asn	CAT His	TAC Tyr	AAC Asn 110	ACC Thr	TAT Tyr	ATA Ile	TCC Ser	AAG Lys 115	AAG Lys	CAT His	GCA Ala	GAG Glu	AAG Lys 120	450
	AAT Asn	TGG Trp	TTT Phe	GTT Val	GGC Gly 125	CTC Leu	AAG Lys	AAG Lys	AAT Asn	GGG Gly 130	AGC Ser	TGC Cys	AAA Lys	CGC Arg	GGT Gly 135	CCT Pro	498
	CGG Arg	ACT Thr	CAC His	TAT Tyr 140	GGC Gly	CAG Gln	AAA Lys	GCA Ala	ATC Ile 145	TTG Leu	TTT Phe	CTC Leu	CCC Pro	CTG Leu 150	CCA Pro	GTC Val	546
		TCT Ser			AGAG	ATC '	TGTT	CTGG	GT G	TTGA	CCAC	T CC	AGAG	AAGT	TTC	GAGGGG	604

# TCCTCACCTG GTTGACCCAA AAATGTTCCC TTGA

638

60 103

# (2) INFORMATION FOR SEQ ID NO:4:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 155 amino acids
  - (B) TYPE: amino acid
  - (C) STRANDEDNESS: single
  - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: protein
- (v) FRAGMENT TYPE: internal
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

Met Ala Glu Gly Glu Ile Thr Thr Phe Thr Ala Leu Thr Glu Lys Phe Asn Leu Pro Pro Gly Asn Tyr Lys Lys Pro Lys Leu Leu Tyr Cys Ser 25 Asn Gly Gly His Phe Leu Arg Ile Leu Pro Asp Gly Thr Val Asp Gly 40 Thr Arg Asp Arg Ser Asp Gln His Ile Gln Leu Gln Leu Ser Ala Glu Ser Val Gly Glu Val Tyr Ile Lys Ser Thr Glu Thr Gly Gln Tyr Leu 75 Ala Met Asp Thr Asp Gly Leu Leu Tyr Gly Ser Gln Thr Pro Asn Glu 90 85 Glu Cys Leu Phe Leu Glu Arg Leu Glu Glu Asn His Tyr Asn Thr Tyr 105 Ile Ser Lys Lys His Ala Glu Lys Asn Trp Phe Val Gly Leu Lys Lys 120 Asn Gly Ser Cys Lys Arg Gly Pro Arg Thr His Tyr Gly Gln Lys Ala 135

### (2) INFORMATION FOR SEQ ID NO:5:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 103 base pairs

Ile Leu Phe Leu Pro Leu Pro Val Ser Ser Asp

- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:

AGAATTCAAA TATTCTGAAA TGAGCTGTTG ACAATTAATC ATCGAACTAG TTAACTAGTA CGCAAGTTCA CGTAAAAAAGG GTATCACATA TGGTACCTGC AGA

(2) INFORMATION FOR SEQ ID NO:6:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 103 base pains
  - (B) TYPE: nucleic acid
  - (C) STRANDEDNESS: single
  - (D) TOPOLOGY: linear

1	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:	
TCTGCAGGTA CCATATGTGA TACCCTTTTT ACGTGAACTT GCGTACTAGT TAACTAGTTC GATGATTAAT TGTCAACAGC TCATTTCAGA ATATTTGAAT TCT	60 103
(2) INFORMATION FOR SEQ ID NO:7:	
(i) SEQUENCE CHARACTERISTICS:	
(A) LENGTH: 16 base pairs	
(B) TYPE: nucleic acid	
(C) STRANDEDNESS: single (D) TOPOLOGY: linear	
~	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:	
	16
GAAATACACC AGTTGG	10
(2) INFORMATION FOR SEQ ID NO:8:	
(i) SEQUENCE CHARACTERISTICS:	
(A) LENGTH: 17 base pairs	
(B) TYPE: nucleic acid	
(C) STRANDEDNESS: single	
(D) TOPOLOGY: linear	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:	
(XI) SEQUENCE DESCRIPTION. SEQ IS NO. 8.	
ACTTGGATCC AAAACAG	17
CONTRACTON FOR CEO TO MO. C.	
(2) INFORMATION FOR SEQ ID NO:9:	
(i) SEQUENCE CHARACTERISTICS:	
(A) LENGTH: 25 base pairs	
(B) TYPE: nucleic acid	
(C) STRANDEDNESS: single	
(D) TOPOLOGY: linear	
! :	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:	
TTTTACATGA AGCTTTATAT TTCAG	25
(2) INFORMATION FOR SEQ ID NO:10:	
(i) SEQUENCE CHARACTERISTICS:	
(A) LENGTH: 32 base pairs	
(B) TYPE: nucleic acid	
(C) STRANDEDNESS: single	
(D) TOPOLOGY: linear	

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:	
AGCTTCATAT GGCTGCTGGT TCTATCACTA CC	32
(2) INFORMATION FOR SEQ ID NO:11:	
(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 28 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:	
CTGCCAGCTC TGCCAGAAGA CGGTGGTT	28
(2) INFORMATION FOR SEQ ID NO:12:	
(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 28 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:12:	
CTGGTGCCTT CCCACCAGGT CACTTCAA	28
(2) INFORMATION FOR SEQ ID NO:13:	
(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 28 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:13:	
AGACCCAAAA CGTCTGTACT GCAAAAAC	28
(2) INFORMATION FOR SEQ ID NO:14:  (i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 19 base pairs  (B) TYPE: nucleic abid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:14:	

GGTGGTTTCT TCCTGCGCA

(2) INFORMATION FOR SEQ ID NO:15:	
(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:15:	
TAGAACCAGC AGCCA†ATGA	20
(2) INFORMATION FOR SEQ ID NO:16:	
(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 28 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear  (xi) SEQUENCE DESCRIPTION: SEQ ID NO:16:  TCTTCTGGCA GAGCTGGCAG GGTAGTGA	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:16:	
TCTTCTGGCA GAGCTGGCAG GGTAGTGA	28
(2) INFORMATION FOR SEQ ID NO:17:	
(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 28 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:17:	
ACCTGGTGGG AAGGCACCAG AACCACCG	28
(2) INFORMATION FOR SEQ ID NO:18:	
(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 28 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:18:	
	28
AGTACAGACG TTTTGGGTCT TTGAAGTG	20
(2) INFORMATION FOR SEQ ID NO:19:	
<ul><li>(i) SEQUENCE CHARACTERISTICS:</li><li>(A) LENGTH: 31 base pairs</li><li>(B) TYPE: nucleic acid\(\)</li></ul>	

(C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(b) Torollog1. IThear	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:19:	
AGCTTGCGCA GGAAGAACC ACCGTTTTTG C	31
(2) INFORMATION FOR SEQ ID NO:20:	
(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 135 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:20:	
AGCTTCATAT GGCTGCTGGT TCTATCACTA CCCTGCCAGC TCTGCCAGAA GACGGTGGTT CTGTGCCTT CCCACCAGGT CACTTCAAAG ACCCAAAACG TCTGTACTGC AAAAACGGTG GTTTCTTCCT GCGCA	60 120 135
(2) INFORMATION FOR SEQ ID NO:21:	
(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 135 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:21:	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:21:  AGCTTGCGCA GGAAGAAACC ACCGTTTTTG CAGTACAGAC GTTTTGGGTC TTTGAAGTGA CCTGGTGGGA AGGCACCAGA ACCACCGTCT TCTGGCAGAG CTGGCAGGGT AGTGATAGAA CCAGCAGCCA TATGA	60 120 135
(2) INFORMATION FOR SEQ ID NO:22:	
(i) SEQUENCE CHARACTER STICS:  (A) LENGTH: 23 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:22:	
ACGTCTGTAC TCCAAAAACG GTG	23
(2) INFORMATION FOR SEQ ID NO:23:	
(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 23 base pairs  (B) TYPE: nucleic acid	
(C) STRANDEDNESS: single	

(D) TOPOLOGY: linear	
A TO NO 22	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:23:	
TACAGACGAG TCTTTTTTTTG	23
(2) INFORMATION FOR SEQ ID NO:24:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 23 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> <li>(xi) SEQUENCE DESCRIPTION: SEQ ID NO:24:</li> </ul>	
CAAAGGAGTG TCTGCAAACC GTT	23
(2) INFORMATION FOR SEQ ID NO:25:	
(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 23 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
A STORES TO NO. 25	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:25:	
AGCTTCTAAA TCTGTTACAG ACG	23
(2) INFORMATION FOR SEQ ID NO:26:	
(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 23 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:26:	
ACGTCTGTAC TCCAAAAACG GTG	23
(2) INFORMATION FOR SEQ ID NO:27:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 23 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	

	1	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:27:	
	TACAGACGAG TCTTTCTTTT TTG	23
	(2) INFORMATION FOR SEQ ID NO:28:	
	(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 24 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:28:	
	GTCTGTACTG CTCAAACGGT GGTT	24
,	(2) INFORMATION FOR SEQ ID NO:29:	
, ,	(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 23 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:29:	
	TTTCTTCCTG CTCATCCACC CCG	23
	(2) INFORMATION FOR SEQ ID NO:30:  (i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 23 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:30:	2.2
	CATCCACCCC GCCGGCCGAG TGG	23
	(2) INFORMATION FOR SEQ ID NO:31:  (i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 23 base pairs  (B) TYPE: nucleid acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:31:	
	CCCCGACGGC CTAGTGGACG GGG \	23

(2) INFORMATION FOR SEQ ID NO:32:	
(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 24 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:32:	
	24
ACCCCGACGG CGCACTGGAC GGGG	2.3
(2) INFORMATION FOR SEQ ID NO:33:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 24 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:33:	
CGCCGGAGT GGCCGGGGTC CGCG	24
(2) INFORMATION FOR SEQ ID NO:34:	
CGGCCGGAGT GGCCGGGGTC CGCG  (2) INFORMATION FOR SEQ ID NO:34:  (i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 25 base pairs  (B) TYPE: nupleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear  (xi) SEQUENCE DESCRIPTION: SEQ ID NO:34:	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:34:	
GAGTGGACGG GAAACGCGAG AAGAG	25
(2) INFORMATION FOR SEQ ID NO:35:	
(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 23 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:35:	
GGACGGGGTC CTCGAGAAGA GCG	23
(2) INFORMATION FOR SEQ ID NO:36:	
<ul><li>(i) SEQUENCE CHARACTERISTICS:</li><li>(A) LENGTH: 23 base pairs</li><li>(B) TYPE: nucleic acid</li></ul>	

	(C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(b) Tolodosi. Ilmodi	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:36:	
	GGTCCGCGAG ATGAGGGACC CAC	23
	(2) INFORMATION FOR SEQ ID NO:37:	
	(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 24 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:37:	
	GGTCCGCGAG ATAAGCGACC CACA	24
	(2) INFORMATION FOR SEQ ID NO:38:	
	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 23 base pairs (B) TYPE: nucle↓c acid	
	(C) STRANDEDNESS: single	
	(C) STRANDEDNESS, STIGTE (D) TOPOLOGY: lihear	
	(b) TOPOLOGI. II lear	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:38:	
	CGAGAAGAGC GCCCCACACA TCA	23
	(2) INFORMATION FOR SEQ ID NO:39:	
	(i) SEQUENCE CHARACTERISTICS:	
·.,	(A) LENGTH: 23 base pairs	
1	(B) TYPE: nucleic acid	
/	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:39:	
	GAGCGACCCA AACATCAAAC TAC	23
	(2) INFORMATION FOR SEQ ID NO:40:	
	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 24 base pairs	
	(B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	

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(xi) SEQUENCE DESCRIPTION: SEQ ID NO:40:	
AGAAGATGGA ACTTTACTAG CTTC	24
(2) INFORMATION FOR SEQ ID NO:41:	
(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 23 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	•
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:41:	
ATGTGTTACA GCAGAGTGTT TCT	23
(2) INFORMATION FOR SEQ ID NO:42:	
(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH 25 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:42:	
GTTACAGACG CCTGTTTCTT TTTTG	25
(2) INFORMATION FOR SEQ ID NO:43:	
(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 25 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:43:	
GTGTTACAGA CAGTTGTTTC TTTTT	25
(2) INFORMATION FOR SEQ ID NO:44:	
(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 24 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:44:	
CTTTCTTTT TTCACCATTC CAC $\dot{\mathbf{T}}$	2.4

(2) INFORMATION FOR SEQ ID NO: 45:	
(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 23 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:45:	
CTTTTTTGAA CTATTGGAGT CTA	23
(2) INFORMATION FOR SEQ ID NO:46:	
(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 24 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:46:	
TGAACGATTG GCATCTAATA ACTA	24
(2) INFORMATION FOR SEQ ID NO:47:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 25 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:47:	
AGTCTAATAA CGCAAATACT TACCG	<b>2</b> 5
(2) INFORMATION FOR SEQ ID NO:48:	
(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 23 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESB: single  (D) TOPOLOGY: linear	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:48:	
CTAATAACTA CAGTACTTAC CGG	23
(2) INFORMATION FOR SEQ ID NO:49:	
<ul><li>(i) SEQUENCE CHARACTERISTICS:</li><li>(A) LENGTH: 24 base pairs</li><li>(B) TYPE: nucleic acid</li></ul>	

(C) \$TRANDEDNESS: single (D) TOPOLOGY: linear	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:49:	
CAATACTTAC ACTTCAAGGA AATA	24
(2) INTORMATION FOR SEQ ID NO:50:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 23 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	
(xi) SEQUENÇE DESCRIPTION: SEQ ID NO:50:	
CAATACTTAC CTGTCAAGGA AAT	23
(2) INFORMATION FOR SEQ ID NO:51:	
(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 25 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear  (xi) SEQUENCE DESCRIPTION: SEQ ID NO:51:	
ACCGGTCAAG GTCTTACACC AGTTG	25
(2) INFORMATION FOR SEQ ID NO:52:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 24 base pairs</li> <li>(B) TYPE: nucleit acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:52:	
GGTGGTTTCT TCCACCCCGA CGGC	24
(2) INFORMATION FOR SEQ ID NO:53:	
(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 24 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear;	

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:53:	
CCCGACGGCC GAGTCCGCGA GAAG	24
(2) INFORMATION FOR SEQ ID NO:54:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 24 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:54:	
CCACACATCA AACAAGCAGA AGAG	24
(2) INFORMATION FOR SEQ ID NO:55:	
(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 24 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:55:	
GCAAACCGTT ACAAAGAAGA TGGA	24
(2) INFORMATION FOR SEQ ID NO:56:	
(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 24 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY linear	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:56:	
TGTTTCTTTT TTGAGTCTAA TAAC	24
(2) INFORMATION FOR SEQ ID NO:57:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 24 pase pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:57:	
GAGTCTAATA ACTACCGGTC AAGG	24

(2) INFORMATION FOR SEQ ID NO:58:	
(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 24 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:58:	
AGGTCACTTC ATGGACCCAA AACG	24
(2) INFORMATION FOR SEQ ID NO:59:	
(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 24 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: \linear	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:59:	
TCAAAGACCC AGCACGTCTG TACT	24
(2) INFORMATION FOR SEQ ID NO:60:	
(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 24 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:60:	
AAGACCCAAA ATCTCTGTAC TGCA	24
(2) INFORMATION FOR SEQ ID NO:61:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 25 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:61:	
GTCACTTCAA AAAGCCAAAA CGTCT	25
(2) INFORMATION FOR SEQ ID NO:62:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 23 base pairs (B) TYPE: nucleic acid	

	(C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:62:	
CAATAC	TTAC CTGTÇAAGGA AAT	23
	(2) INFORMATION FOR SEQ ID NO:63:	
	<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 24 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:63:	
GTCTGT	ACTG CTCAAACGGT GGTT	24
	(2) INFORMATION FOR SEQ ID NO:64:	
	<ul><li>(i) SEQUENCE CHARACTERISTICS:</li><li>(A) LENGTH: 25 base pairs</li><li>(B) TYPE: nucleic acid</li><li>(C) STRANDEDNESS: single</li><li>(D) TOPOLOGY: linear</li></ul>	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:64:	
ATGTGG	GCACT GTCTCGAACT GGGCA	25
	(2) INFORMATION FOR SEQ ID NO:65:	
	<ul><li>(i) SEQUENCE CHARACTERISTICS:</li><li>(A) LENGTH: 25 base pairs</li><li>(B) TYPE: nucleic acid</li><li>(C) STRANDEDNESS: single</li><li>(D) TOPOLOGY: linear</li></ul>	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:65:	
ATGTGG	GCACT GGAGCGAACT GGGCA	25
	(2) INFORMATION FOR SEQ ID NO:66:	
	<ul><li>(i) SEQUENCE CHARACTERISTICS:</li><li>(A) LENGTH: 23 base pairs</li><li>(B) TYPE: nucleic acid</li><li>(C) STRANDEDNESS: single</li><li>(D) TOPOLOGY: linear</li></ul>	

	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:66:	
GGCAC	TGAAA ACTACTGGGC AGT	23
	(2) INFORMATION FOR SEQ ID NO:67:	
	<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 25 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:67:	
CTGGG	CAGTA TTCTCTTGGA TCCAA	25
	(2) INFORMATION FOR SEQ ID NO:68:  (i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 25 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear  (xi) SEQUENCE DESCRIPTION: SEQ ID NO:68:	
AACTT	GGATC CTCTACAGGA CCTGG	25
	(2) INFORMATION FOR SEQ ID NO:69:  (i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 36 base/pairs  (B) TYPE: nucleic adid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear  (ii) MOLECULE TYPE: cDNA  (xi) SEQUENCE DESCRIPTION: SEQ ID NO:69:	
TTGGG	TCTTT GAAGTGCATA TGTGGGAAGG CACCAG	36